FORM PTO-1449 (MODIFIED) Applicant(s): Rice et al. Docket No.: YOR920030512US1 LIST OF PUBLICATIONS FOR Serial No.: 10/699,373 APPLICANT'S INFORMATION Filing Date: October 31, 2003 DISCLOSURE STATEMENT Group: TENT DOCUMENTS FILING DATE EVAMINED CLASS/SUBCLASS IF APPROPRIATE INITIAL FOREIGN PATENT DOCUMENTS TRANSLATION EXAMINER INITIAL DOCUMENT NO. DATE COUNTRY CLASS/SUBCLASS YES NO OTHER DOCUMENTS EXAMINER INITIAL REF NO. AUTHOR, TITLE, DATE, PERTINENT PAGES, ETC. Akutsu et al., "Algorithms for Inferring Qualitative Models of Biological Networks," Proc. Pacific Symposium on Biocomputing 2000 (PSB'2000), pp. 290-301 (2000). Friedman et al., "Using Bayesian Networks to Analyze Expression Data," Journal of Computational Biology, Vol. 7, Nos. 3/4, pp. 601-620 (2000). Hartemink et al., "Using Graphical Models and Genomic Expression Data to Statistically Validate Models of Genetic Regulatory Networks," Pacific Symposium on Biocomputing 2001, Hawaii, January 2001. D'haeseleer et al., "Genetic Network Inference: From Co-Expression Clustering to Reverse Engineering," Bioinformatics, Vol. 16, No. 8 pp. 707-726 (2000). Samoilov et al., "On the Deduction of Chemical Reaction Pathways from Measurements of Time Series of Concentrations," CHAOS, vol. 11, no. 1, pp. 108-114 (2001). Arkin et al., "A Test Case of Correlation Metric Construction of a Reaction Pathway from Measurements," Science, Vol. 277, no. 29 pp. 1275-1279 (1997). Arkin et al., "Statistical Construction of Chemical Reaction Mechanisms from Measured Time-Series," J. Phys. Chem., vol. 99, pp. 970-79 (1995). A. Wagner, "Estimating Coarse Gene Network Structure from Large-Scale Gene Perturbation Data," Genome Research, vol. 12, pp. 309-315 (2002).

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